HMO Glycan Microarray User Manual



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Introduction

Human milk oligosaccharides (HMOs) are the third most abundant solid ingredient in breast milk. These sugars - unique to human milk - protect newborns from disease and infection, establish healthy gut bacteria, and are essential for infant brain, intestinal, and immune system development.

Inaugural studies of HMOs have found many potential benefits for the incorporation of HMOs in infant formula as well as medications for adults. They have been found not only to restore beneficial microorganisms to the gut, but also function as a natural antiviral, inhibit growth of pathogenic bacteria, and inhibit tumor cell growth. In addition, HMO profiling has been used to evaluate infant digestive development, diarrhea incidence, morbidity and mortality, body composition, and risk of developing certain disorders.

Upon discovering these highly beneficial roles for HMOs, biopharmaceutical companies have begun to incorporate 1 or 2 of the abundant HMOs in their infant formula, and it is anticipated that more HMOs will be added in the future to better mimic the complete nutrition provided by natural breastmilk. However, the specific functions and interactions of these carbohydrates are still largely unstudied and there remains much potential for their application as biomarkers or in the development of drugs and infant formula.

Z Biotech's HMO Glycan Microarray is designed to make the investigation of HMOs easy and efficient. As leaders in the glycan microarray industry we have developed sensitive, high density arrays for the research of glycan-binding proteins and antibodies. Using minimal sample volume and only a few hours, these arrays can be tested with cells, antibodies, or other proteins to determine which HMOs interact and give insight into the functions of these HMOs and their binding partners.

This manual is provided as a comprehensive guide to help the researcher acquire clear results from the assay. Please read through carefully before starting your experiment.

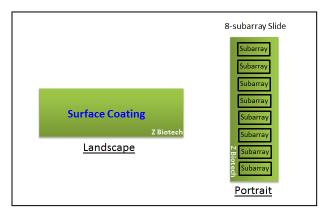
Handling and Storage

Store the bag of slides and any buffers in a 4°C refrigerator if they are to be assayed within 3 weeks upon receipt. For long term storage keep the bag of slides at -20°C. Avoid freezing and thawing multiple times. Purchased slides and buffers should be used within 6 months.

Allow the bag of slides to equilibrate to room temperature at least 20 minutes before opening. After opening, re-seal any unused slides in the moisture barrier bag with a desiccant inside and refreeze. Handle the slides in a clean, dust free environment. Wear gloves and hold the slides on the edges. When adding sample do not touch the pipette tip to the array surface. When removing sample gently touch the pipette tip at the corner of the well of the cassette and tip the slide so that the sample pools to that corner. Avoid contact with the surface of the slides.

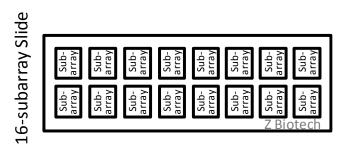
Array Map/Schematic

HMO Glycan Microarray slides have 8 or 16 subarrays depending on the customer's request. Arrays are printed on the side with the "Z Biotech" label facing upward. The "Z Biotech" label is located on the bottom right corner from a landscape point of view, or the bottom left corner in a portrait point of view (see image on right). The orientation of the printed array is consistent with the portrait slide orientation. Dimensions and array map are shown on the following pages.



Array Map (16-subarray):

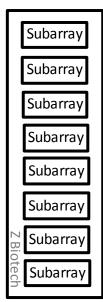
	1	2	3	4	5	6	7	8	9	10	11	12
1	HMO1	HMO1	HMO1	HMO2	HMO2	HMO2	нмоз	нмоз	нмоз	HMO4	HMO4	HMO4
2	HMO5	HMO5	HMO5	нмо6	нмо6	нмо6	нмо7	НМО7	нмо7	HM08	HMO8	HMO8
3	нмо9	нмо9	нмо9	HMO10	HMO10	HMO10	HM011	HMO11	HMO11	HMO12	HMO12	HMO12
4	HMO13	HMO13	HMO13	HMO14	HMO14	HMO14	HM015	HMO15	HMO15	HM016	HMO16	HMO16
5	HM017	HMO17	HM017	HMO18	HMO18	HMO18	HMO19	HMO19	HMO19	HMO20	HMO20	HMO20
6	HMO21	HMO21	HMO21	HMO22	HMO22	HMO22	HMO23	HMO23	HMO23	HMO24	HMO24	HMO24
7	HMO25	HMO25	HMO25	HMO26	HMO26	HMO26	HMO27	HMO27	HMO27	HMO28	HMO28	HMO28
8	HMO29	HMO29	HMO29	НМО30	нмозо	нмозо	HMO31	HMO31	HMO31	HMO32	HMO32	HMO32
9	НМО33	HMO33	НМО33	HMO34	HMO34	HMO34	HMO35	HMO35	HMO35	нмоз6	нмоз6	нмоз6
10	HM037	HMO37	HMO37	HMO38	НМО38	НМО38	нмоз9	нмоз9	НМО39	HMO40	HMO40	HMO40
11	HM041	HMO41	HM041	HMO42	HMO42	HMO42	HMO43	HMO43	HMO43	HM044	HMO44	HMO44
12	HM045	HM045	HM045	HM046	HM046	HM046	NC	NC	NC	PC1	PC1	PC1
13	PC2	PC2	PC2	PC3	PC3	PC3	PC4	PC4	PC4	MARKER	MARKER	MARKER



Array Map (8-subarray):

25 26 27 28 29	нмоз нмоз нмоз нмоз	нмото нмото нмото нмото	HMO15 HMO15 HMO15 HMO15	нмого нмого нмого нмого	нмогь нмогь нмогь нмогь	нмозо нмозо нмозо нмозо	нмозь нмозь нмозь нмозь	HMO40 HMO40 HMO40 HMO40	HMO45 HMO45 HMO45 HMO45 HMO45	PG PG PG PC3 PC3	
23 24	HMO4 HMO4 HI	ин 60мн 60мн	HM014 HM014	нмоте нмоте	HM024 HM024	НМО29 НМО29	HM034 HM034	нмоз9 нмоз9	HMO44 HMO44	PC2 PC2	
20 21 22	4 HMO4 HMO4 HMO4	60МН 60МН 6	4 HM014 HM014 HM014	9 HM019 HM019 HM019	4 HMO24 HMO24 HMO24	9 HMO29 HMO29	4 HMO34 HMO34 HMO34	9 HMO39 HMO39	4 HMO44 HMO44 HMO44	PC2 PC2 PC2	
17 18 19	нмоз нмоз нмо4	нмов нмов	нмо13 нмо13 нмо14	нмотв нмотв нмотя	нмогз нмогз нмог4	нмо28 нмо28 нмо29	нмозз нмозз нмоз4	нмозв нмозв нмоз9	HMO43 HMO44	PCI PCI PCZ	
15 16	нмоз нмоз н	н н н н н н н н н н н	нмо13 нмо13 н	нмотв нмотв н	нмогз нмогз н	нмо28 нмо28 н	нмозз нмозз ні	нмозв нмозв ні	нмо43 нмо43 н	PC1 PC1	
3 14	HM03	HM08	HM013	HM018	HM023	HM028	нмозз	HM038	HM043	1 PCI	
12 13	2 нмоз нмоз	7 HM07 HM08	12 HM012 HM013	17 HM017 HM018	22 HM022 HM023	27 HM027 HM028	32 HMO32 HMO33	37 HMO37 HMO38	12 HMO42 HMO43	NC PCI	
10 11	нмо2 нмо2	нмо7 нмо7	HM012 HM012	HMO17 HMO17	HM022 HM022	HMO27 HM027	нмоз2 нмоз2	HM037 HM037	HM042 HM042	NC	
6	эг нмог нмог	70MH TOMH TC	112 HMO12 HMO12	117 HMO17 HMO17	122 HMO22 HMO22	727 HMO27 HMO27	332 НМО32 НМО32	337 HMO37 HMO37	342 HMO42 HMO42	NC NC	
9	1 HMO1 HMO2	нмое нмо7	11 HM011 HM012	16 HM016 HM017	21 HMO21 HMO22	26 HMO26 HMO27	31 HMO31 HMO32	36 HMO36 HMO37	41 HMO41 HMO42	46 HMO46 NC	
4 5	11 HM01 HM01	90MH 90MH 90	11 HM011 HM011	нмоте нмоте нмоте	21 HMO21 HMO21	26 нмо26 нмо26	31 HMO31 HMO31	нмозе нмозе нмозе	HMO41 HMO41 HMO41	HM046 HM046 HM046	
2 3	нмот нмот	нмое нмое	HM011 HM011	HM016	HM021 HM021	нмо26 нмо26	HM031 HM031	НМОЗ6	HMO41	HMO46	
1	1 HM01	2 HM06	3 HM011	4 HM016	5 HM021	9 нмозе	7 HM031	8 HM036	9 нмо41	10 HM046	

8-subarray Slide



HMO Identification List:

ID	Structure	Common Name					
HMO1	Galβ1-4Glc	Lactose					
HMO2	Galβ1-4(Fucα1-3)Glc	3-FL					
HMO3	Fucα1-2Galβ1-4Glc	2'-FL					
HMO4	Neu5Acα2-3Galβ1-4Glc	3'-SL					
HMO5	Neu5Acα2-6Galβ1-4Glc	6'-SL					
HMO6	Galα1-4Galβ1-4Glc	Gb ₃ (P ^k antigen)					
HMO7	GalNAcβ1-3Galα1-4Galβ1-4Glc	Gb ₄					
HMO8	Galβ1-3GalNAcβ1-3Galα1-4Galβ1-4Glc	Gb ₅ (SSEA-3)					
HMO9	Fucα1-2Galβ1-3GalNAcβ1-3Galα1-4Galβ1-4Glc	Globo-H					
HMO10	Galβ1-3GlcNAcβ1-3Galβ1-4Glc	LNT					
HMO11	Galβ1-4GlcNAcβ1-3Galβ1-4Glc	LNnT					
HMO12	Fucα1-2Galβ1-3GlcNAcβ1-3Galβ1-4Glc	LNFP-I					
HMO13	Galβ1-3(Fucα1-4)GlcNAcβ1-3Galβ1-4Glc	LNFP-II					
HMO14	Galβ1-4(Fucα1-3)GlcNAcβ1-3Galβ1-4Glc	LNFP-III					
HMO15	Fucα1-2Galβ1-4GlcNAcβ1-3Galβ1-4Glc	Blood group H antigen					
HMO16	Fucα1-2Galβ1-4(Fucα1-3)GlcNAcβ1-3Galβ1-4Glc	LeY					
HMO17	Neu5Acα2-6Galβ1-4GlcNAcβ1-3Galβ1-4Glc	LsTc-I					
HMO18	Galβ1-4GlcNAcβ1-3(Neu5Acα2-6)Galβ1-4Glc	LsTc-II					
HMO19	Neu5Acα2-3Galβ1-3GlcNAcβ1-3Galβ1-4Glc	LsTa					
HMO20	Neu5Acα2-3Galβ1-4GlcNAcβ1-3Galβ1-4Glc	Ac3-LNnT					
HMO21	Neu5Acα2-6Galβ1-4GlcNAcβ1-3Galβ1-4Glc	Ac6-LNnT					
HMO22	Neu5Acα2-3Galβ1-3(Neu5Acα2-6)GlcNAcβ1-3Galβ1-4Glc	DSLNT					
HMO23	GlcNAcβ1-3Galβ1-4GlcNAcβ1-3Galβ1-4Glc	Lacto-N-neoPentaose (LNnP)					
HMO24	GlcNAcβ1-3Galβ1-4(Fucα1-3)GlcNAcβ1-3Galβ1-4Glc	Fuc-LNnP					
HMO25	GlcNAcβ1-6(GlcNAcβ1-3)Galβ1-4Glc	N/A					
HMO26	GlcNAcβ1-6(Galβ1-4GlcNAcβ1-3)Galβ1-4Glc	N/A					
HMO27	GlcNAcβ1-6(Galβ1-3GlcNAcβ1-3)Galβ1-4Glc	N/A					
HMO28	GlcNAcβ1-6(Neu5Acα2-6Galβ1-4GlcNAcβ1-3)Galβ1-4Glc	N/A					
HMO29	GlcNAcβ1-6(Neu5Acα2-6Galβ1-3GlcNAcβ1-3)Galβ1-4Glc	N/A					
HMO30	GlcNAcβ1-6(Neu5Acα2-3Galβ1-3GlcNAcβ1-3)Galβ1-4Glc	N/A					
HMO31	GlcNAcβ1-6(Galβ1-4(Fucα1-3)GlcNAcβ1-3)Galβ1-4Glc	N/A					
HMO32	GlcNAcβ1-6(Fucα1-2Galβ1-4GlcNAcβ1-3)Galβ1-4Glc	N/A					
HMO33	GlcNAcβ1-6(Fucα1-2Galβ1-4(Fucα1-3)GlcNAcβ1-3)Galβ1-4Glc	N/A					
HMO34	Galβ1-4GlcNAcβ1-6(GlcNAcβ1-3)Galβ1-4Glc	N/A					
HMO35	Neu5Acα2-6Galβ1-4GlcNAcβ1-6(GlcNAcβ1-3)Galβ1-4Glc	N/A					
HMO36	Galβ1-4(Fucα1-3)GlcNAcβ1-6(GlcNAcβ1-3)Galβ1-4Glc	N/A					
HMO37	Fucα1-2Galβ1-4GlcNAcβ1-6(GlcNAcβ1-3)Galβ1-4Glc	N/A					
HMO38	Galβ1-4GlcNAcβ1-6(Neu5Acα2-6Galβ1-4GlcNAcβ1-3)Galβ1-4Glc	N/A					
HMO39	Galβ1-4GlcNAcβ1-6(Neu5Acα2-6Galβ1-3GlcNAcβ1-3)Galβ1-4Glc	N/A					
HMO40	Galβ1-4GlcNAcβ1-6(Fucα1-2Galβ1-4GlcNAcβ1-3)Galβ1-4Glc	N/A					
HMO41	Neu5Acα2-6Galβ1-4GlcNAcβ1-6(Galβ1-4GlcNAcβ1-3)Galβ1-4Glc	N/A					
HMO42	Neu5Gcα2-6Galβ1-4GlcNAcβ1-6(Galβ1-4GlcNAcβ1-3)Galβ1-4Glc	N/A					
HMO43	Galβ1-4(Fucα1-3)GlcNAcβ1-6(Galβ1-4GlcNAcβ1-3)Galβ1-4Glc	N/A					
HMO44	Neu5Gcα2-6Galβ1-4GlcNAcβ1-6(Neu5Gcα2-6Galβ1-4GlcNAcβ1-3)Galβ1-4Glc	N/A					
HMO45	Fucα1-2Galβ1-4GlcNAcβ1-6(Fucα1-2Galβ1-4GlcNAcβ1-3)Galβ1-4Glc	N/A					
HMO46	Fucα1-2Galβ1-4(Fucα1-3)GlcNAcβ1-6(Fucα1-2Galβ1-4(Fucα1-3)GlcNAcβ1-3)Galβ1-4Glc	N/A					

Controls

NC: Negative control, Print Buffer

PC1: Positive control 1, a biotinylated probe (0.01 mg/ml)

PC2: Postitive control 2, Human IgG (0.1 mg/ml)

PC3: Postitive control 3, Mouse IgG (0.1 mg/ml)

PC4: Postitive control 4, Rabbit IgG (0.1 mg/ml)

Array Marker: Anti-Human IgG, Cy3 (0.01 mg/ml) and anti-Human IgG, Alexa555 (0.01 mg/ml)

Materials Required

- Arrayed glass slide
- 16 or 8 subarray cassette
- Blocking Buffer (Item #10106)
- Glycan Array Assay Buffer (GAAB Item #10107)

Materials above are available for purchase

- Wash Buffer: 20 mM Tris-HCl, 150 mM NaCl, 0.05% Tween 20, pH 7.6
- Glycan-binding protein samples of interest
- Biotinylated secondary antibodies (for sandwich assay format only)
- Fluorescent labeled streptavidin (for biotin labeled glycan-binding samples)
- Sterile de-ionized water
- Orbital shaker
- Laser fluorescence scanner (able to scan at the wavelength of your fluorophore)
- Centrifuge
- Pipette and sterile pipette tips
- Sterile centrifuge tubes
- Coplin jar or 250 mL beaker
- Adhesive slide cover film
- Aluminum foil

Preparation of assay samples:

Prepare glycan-binding samples or detection antibodies in a centrifuge tube by diluting with the GAAB buffer. For the fluorescently labelled streptavidin we recommend a concentration of 1 μ g/mL. For detection antibodies, we recommend a concentration around 1-10 μ g/ml. We generally recommend a range of 100 μ g/ml to 0.1 μ g/ml concentration for glycan-binding samples, although some experimentation may be required to establish the concentration that will provide the highest binding signals with the lowest background fluorescence. This is often accomplished by applying a different dilution of samples to different wells of the array. In addition to testing a dilution range for your glycan-binding protein of interest, we recommend setting up control assays for any additional detection or secondary antibodies to ensure that any binding observed is specific to your protein of interest. Fluorescent signal due to specific binding to your protein of interest should be dose-dependent within the dynamic range of your protein dilution, and should have positive binding signal after signal from control assays has been subtracted. Calculate the volume

of sample needed depending on how many slides and subarrays are to be assayed. We recommend using 100 μ L volume of sample per well for 16 subarray cassettes and 200 μ L for 8 subarray cassettes to ensure full and even coverage of the printed area throughout incubation. If necessary, the assay can be done successfully with a minimal volume of 60 μ L per well for 16 subarray cassettes and 80 μ L for 8 subarrays. We caution that using a minimal volume in the wells has an increased risk of the array drying out during the assay, and may also cause unequal distribution of the sample across the arrayed surface which may result in signal variation. Please ensure each sample is homogeneous and thoroughly mixed, and there are no dry spots on the subarray after the sample is added.

Assay Protocol

Part 1 – Blocking

Handle the slide in a clean, dry environment. Use gloves and avoid touching the slide surface

- 1. Allow the arrayed slides to equilibrate to room temperature (20-30 minutes) before opening the moisture barrier bag.
- 2. Add blocking buffer to each subarray well. We recommend using 100 μ L per well for 16 subarray cassettes and 200 μ L for 8 subarray cassettes to ensure full and even coverage of the printed area throughout incubation.
- 3. Cover the wells with adhesive film to prevent evaporation and incubate slide on shaker at 85 rpm for 1 hour. Longer incubation time is acceptable, but not necessary.

Make sure the orbital shaker is completely flat. If the slide is sloped in any direction during incubation it can cause variation in binding.

Part 2 – Binding assay

- 1. Unless the glycan-binding sample of interest is bacteria or cells, centrifuge samples briefly to avoid adding irrelevant particles to the array.
- 2. Remove blocking buffer from each well by gently touching the pipette tip to the corner of the well of the cassette and tip the slide so that the sample pools to that corner. Avoid touching the array surface.
- 3. Immediately apply the glycan-binding sample of interest to each well. We recommend using 100 μ L per well for 16 subarray cassettes and 200 μ L for 8 subarray cassettes to ensure full and even coverage of the printed area throughout incubation. Avoid leaving air bubbles.
- 4. Seal the wells with adhesive film to prevent evaporation. If the sample is fluorescently labelled, cover with aluminum foil to keep it in the dark. Incubate on the shaker for 1-3 hours at 100 rpm. If the samples can easily aggregate, shake at higher speed to prevent aggregation. Longer incubation time may increase binding signal, especially for weakly binding samples.

Avoid allowing the slides to dry out at any point during the assay, especially during long incubation times. Make sure the adhesive film is sealed around each well.

If your glycan-binding samples are fluorescently labelled, go directly to Part 6 – Final wash and dry.

Part 3 – Wash

- 1. Remove glycan-binding samples from each well by gently touching the pipette tip to the corner of the well of the cassette and tip the slide so that the sample pools to that corner. Avoid touching the array surface, but a gentle touch is okay to ensure no sample is left pooled in the corners.
- 2. Add wash buffer to each well. We recommend using $100~\mu L$ per well for 16~subarray cassettes and $200~\mu L$ for 8~subarray cassettes. Cover the wells with adhesive film and incubate on the shaker for 5~minutes at 85~rpm. Completely remove the wash buffer by pipette and repeat this step. Avoid allowing the slide to dry out and have your next wash or sample ready before you remove the wash buffer.

If your glycan-binding sample is biotinylated, go directly to Part 5 – Fluorescent staining.

Part 4 – Binding of biotinylated antibody (Sandwich Assay Format)

- 1. Unless the secondary biotinylated antibody sample is bacteria or cells, centrifuge samples briefly to avoid adding irrelevant particles to the array.
- 2. After completely removing the wash buffer immediately add the biotinylated antibody to each well. We recommend using 100 μ L per well for 16 subarray cassettes and 200 μ L for 8 subarray cassettes. Seal the wells with adhesive film and incubate on the shaker for 1 hour at 100 rpm. Longer incubation time is acceptable, but not necessary.
- 3. After incubation repeat Part 3 Wash

Part 5 – Fluorescent staining

- 1. Centrifuge fluorescent-labeled streptavidin samples briefly to avoid adding irrelevant particles to the array.
- 2. After completely removing the wash buffer immediately add the fluorescently labelled streptavidin sample. $100 \, \mu L$ per well is recommended for $16 \, \text{subarray}$ cassettes and $200 \, \mu L$ for 8 subarray cassettes. Seal the wells with adhesive film and shield the wells from light with aluminum foil. Incubate on the shaker at 85 rpm for 1 hour. Longer incubation time is acceptable, but not necessary.

Part 6 – Final wash and dry

- 1. Remove the sample from each well by gently touching the pipette tip to the corner of the well of the cassette and tip the slide so that the sample pools to that corner. Avoid touching the array surface.
- 2. Briefly rinse each well with wash buffer. 100 μ L per well is recommended for 16 subarray cassettes and 200 μ L for 8 subarray cassettes.
- 3. Completely remove the wash buffer by pipette. Avoid touching the array surface. Repeat steps 2 and 3.
- 4. Disassemble the cassette from the slide. For the provided cassette this can be done by holding the slide with one hand at the top and bottom edges and sliding out the cassette clips one by one with the other hand. If your provided cassette has metal clips they can be removed by rotating the clip outwards from the bottom of the slide. When the clips have been removed place the slide on the table and hold a small outer edge of the slide to the table as you gently peel the cassette off.
- 5. Immediately immerse the slide in a coplin jar or beaker full of wash buffer. Do not touch the surface of the array or allow the array surface to touch the sides of the beaker or jar.
- 6. Place the jar or beaker on the 60 rpm shaker for 10 minutes.
- 7. Decant the wash buffer from the jar or beaker while holding the slide in place (only touch the edge of the slide) and then add sterile de-ionized water to immerse the slide.
- 8. Place the jar or beaker on the 60 rpm shaker for 2 minutes.
- 9. Decant the water from the jar or beaker.
- 10. Allow the slide to dry completely in a clean, dust free environment before scanning.

Analysis

Scan the slide in a laser fluorescence scanner at the wavelength of emission for the fluorophore used. Adjust the laser power and PMT to obtain the highest possible signals without any being saturated (saturated positive control signal is okay). Analyze data with microarray analysis software. If there is specific binding the signal intensity should be higher than the background signal (area where there are no printed spots). Fluorescent signal due to specific binding to your sample of interest should be both dose-dependent with your sample dilution (unless the sample concentration range is too high and saturates the spots), and should have positive binding signal after signal from control assays has been subtracted. Our standard method of comparing signal intensities is to quantify the median signal intensity data and subtract the background intensity. Subtracting signal from negative control spots as well as the same spots on a negative control assay (assay with only detection antibodies and fluorophore) will give more accurate specific binding data.

Interpretation of Control Signals:

<u>Negative Control (Print Buffer):</u> The negative control should produce a signal close to the intensity of the background. Since there is no binding involved with the negative control, any other signals around the negative control's intensity are also not binding. Subtracting the negative control's signal from the other binding signals will give more accurate specific binding data.

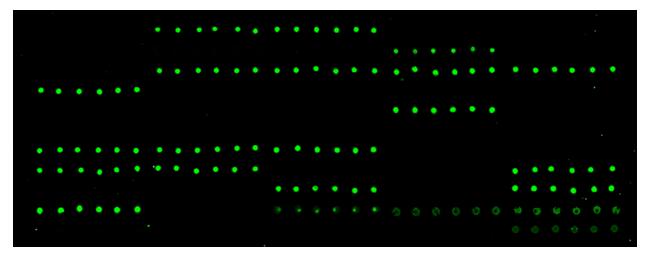
<u>Positive Control (a biotinylated probe):</u> This positive control will bind directly to the fluorescent labelled streptavidin. If your glycan-binding sample is already fluorescently labelled, or in any case where the addition of fluorescent labelled streptavidin to the array was not preformed (Part 5 – Fluorescent staining) this positive control will not be reactive.

<u>IgG (PC2, PC3, PC4)</u>: IgG is an antibody found in blood that is a primary component of humoral immunity. If the glycan-binding or secondary antibody sample is an anti-IgG from human, rabbit, or mouse it should bind to the respective IgG control.

<u>Marker:</u> The array marker should show fluorescence signal regardless of the assay. It is there primarily to aid with orientation of the array map during analysis.

Typical Binding Assay Result from the HMO Glycan Microarray

Example: HMO Glycan Microarray on 8 subarray format. A subarray assayed with biotinylated AAL lectin ($10~\mu g/ml$), followed by streptavidin-Cy3 ($1~\mu g/ml$). The array was scanned with GenePix scanner at 450 PMT and 100% laser power at 532nm wavelength. The positive control shows binding as expected. All HMOs containing fucose show fluorescent binding signal.



Troubleshooting

Condition	Possible Causes	Potential Solutions				
High Background	 Concentration of protein samples is too high. Concentration of fluorescent samples is too high. Arrays are not thoroughly washed. Slide drying out during assay. Excessive particles in the samples due to sample aggregation, dust, etc. 	 Use a lower concentration range of samples. Consider a wider range if you are unsure where the detection limit is. Use control assays to determine which sample is causing high background. Apply longer times for washing steps and use a higher shaking rate Make sure wash buffer and sample is completely removed before the next step. Make sure adhesive film fully seals the wells to avoid evaporation Centrifuge the samples prior to assay to avoid adding irrelevant particles. Make sure buffers are filtered. If you think that the protein is aggregating during incubation, try shaking at a higher speed 				
Signal Variation	 Slide drying out during assay. Binding samples are not equally distributed in the wells Glycan-binding protein aggregation during incubation Bubbles during incubation 	 Make sure wells are sealed to prevent evaporation during incubation. Apply a larger volume of sample to each well to ensure equal distribution Use a higher shaking rate during incubation Make sure samples are homogeneous, mixed thoroughly, and do not leave bubbles on the array surface 				
Unexpected Binding	 Cross contamination between wells or other sources. Sample contamination 	 Make sure to use sterilized pipette tips and tubes used for sample application and preparation Ensure cassette is pressed firmly to the slide so that there are no gaps to allow leaking between wells Be careful not to cross contaminate samples when applying to the wells, even during wash steps 				